## SEQUENCE LISTING

<110> Buchter-Larsen, et al.

<120> A PROCESS OF PREPARING AN ANTI-OXIDANT

<130> 674509-2020

<140> 09/423,126

<141> 1999-11-05

<150> PCT/IB98/00708

<151> 1998-05-06

<150> GB 9709161.5

<151> 1997-05-06

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<170> PatentIn version 3.0

<210> 1

<211> 1088

<212> PRT

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<223> fungus sp. or fungus infected gracilariopsis sp.

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Phe Pro Ala Val His Thr Ala Thr Arg Lys Thr Asn Arg Leu Asn Val 35 40 45

Ser Met Thr Ala Leu Ser Asp Lys Gln Thr Ala Thr Ala Gly Ser Thr 50 55 60

Asp Asn Pro Asp Gly Ile Asp Tyr Lys Thr Tyr Asp Tyr Val Gly Val 65 70 75 80

Trp Gly Phe Ser Pro Leu Ser Asn Thr Asn Trp Phe Ala Ala Gly Ser 85 90 95

Ser Thr Pro Gly Gly Ile Thr Asp Trp Thr Ala Thr Met Asn Val Asn 100 105 110

Phe Asp Arg Ile Asp Asn Pro Ser Ile Thr Val Gln His Pro Val Gln
115 120 125

Val Gln Val Thr Ser Tyr Asn Asn Asn Ser Tyr Arg Val Arg Phe Asn 130 135 140

Pro Asp Gly Pro Ile Arg Asp Val Thr Arg Gly Pro Ile Leu Lys Gln Gln Leu Asp Trp Ile Arg Thr Gln Glu Leu Ser Glu Gly Cys Asp Pro 170 Gly Met Thr Phe Thr Ser Glu Gly Phe Leu Thr Phe Glu Thr Lys Asp Leu Ser Val Ile Ile Tyr Gly Asn Phe Lys Thr Arg Val Thr Arg Lys Ser Asp Gly Lys Val Ile Met Glu Asn Asp Glu Val Gly Thr Ala Ser Ser Gly Asn Lys Cys Arg Gly Leu Met Phe Val Asp Arg Leu Tyr Gly Asn Ala Ile Ala Ser Val Asn Lys Asn Phe Arg Asn Asp Ala Val Lys Gln Glu Gly Phe Tyr Gly Ala Gly Glu Val Asn Cys Lys Tyr Gln Asp Thr Tyr Ile Leu Glu Arg Thr Gly Ile Ala Met Thr Asn Tyr Asn Tyr 280 Asp Asn Leu Asn Tyr Asn Gln Trp Asp Leu Arg Pro Pro His His Asp 290 Gly Ala Leu Asn Pro Asp Tyr Tyr Ile Pro Met Tyr Tyr Ala Ala Pro Trp Leu Ile Val Asn Gly Cys Ala Gly Thr Ser Glu Gln Tyr Ser Tyr Gly Trp Phe Met Asp Asn Val Ser Gln Ser Tyr Met Asn Thr Gly Asp Thr Trp Asn Ser Gly Gln Glu Asp Leu Ala Tyr Met Gly Ala Gln Tyr Gly Pro Phe Asp Gln His Phe Val Tyr Gly Ala Gly Gly Met 370 Glu Cys Val Val Thr Ala Phe Ser Leu Leu Gln Gly Lys Glu Phe Glu 390 Asn Gln Val Leu Asn Lys Arg Ser Val Met Pro Pro Lys Tyr Val Phe 410 Gly Phe Phe Gln Gly Val Phe Gly Thr Ser Ser Leu Leu Arg Ala His 420 Met Pro Ala Gly Glu Asn Asn Ile Ser Val Glu Glu Ile Val Glu Gly 435 440 445

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Tyr	Gln 450	Asn	Asn	Asn	Phe	Pro 455	Phe	Glu	Gly	Leu	Ala 460	Val	Asp	Val	Asp
Met 465	Gln	Asp	Asn	Leu	Arg 470	Val	Phe	Thr	Thr	Lys 475	Gly	Glu	Phe	Trp	Thr 480
Ala	Asn	Arg	Val	Gly 485	Thr	Gly	Gly	Asp	Pro 490	Asn	Asn	Arg	Ser	Val 495	Phe
Glu	Trp	Ala	His 500	Asp	Lys	Gly	Leu	Val 505	Cys	Gln	Thr	Asn	Ile 510	Thr	Cys
Phe	Leu	Arg 515	Asn	Asp	Asn	Glu	Gly 520	Gln	Asp	Tyr	Glu	Val 525	Asn	Gln	Thr
Leu	Arg 530	Glu	Arg	Gln	Leu	Tyr 535	Thr	Lys	Asn	Asp	Ser 540	Leu	Thr	Gly	Thr
Asp 545	Phe	Gly	Met	Thr	Asp 550	Asp	Gly	Pro	Ser	Asp 555	Ala	Tyr	Ile	Gly	His 560
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Gly	Arg	Pro	Asp 580	Val	Ala	Glu	Trp	Trp 585	Gly	Asn	Asn	Tyr	Lys 590	Lys	Leu
Phe	Ser	Ile 595	Gly	Leu	Asp	Phe	Val 600	Trp	Gln	Asp	Met	Thr 605	Val	Pro	Ala
Met	Met 610	Pro	His	Lys	Ile	Gly 615	Asp	Asp	Ile	Asn	Val 620	Lys	Pro	Asp	Gly
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Thr	Tyr	His	Pro	Gln 645	Val	Leu	Val	Thr	Asp 650	Met	Arg	Tyr	Glu	Asn 655	His
Gly	Arg	Glu	Pro 660	Met	Val	Thr	Gln	Arg 665	Asn	Ile	His	Ala	Tyr 670	Thr	Leu
Cys	Glu	Ser 675	Thr	Arg	Lys	Glu	Gly 680	Ile	Val	Glu	Asn	Ala 685	Asp	Thr	Leu
Thr	Lys 690	Phe	Arg	Arg	Ser	Tyr 695	Ile	Ile	Ser	Arg	Gly 700	Gly	Tyr	Ile	Gly
Asn 705	Gln	His	Phe	Gly	Gly 710	Met	Trp	Val	Gly	Asp 715	Asn	Ser	Thr	Thr	Ser 720
Asn	Tyr	Ile	Gln	Met 725	Met	Ile	Ala	Asn	Asn 730	Ile	Asn	Met	Asn	Met 735	Ser
Cys	Leu	Pro	Leu 740	Val	Gly	Ser	Asp	Ile 745	Gly	Gly	Phe	Thr	Ser 750	Tyr	Asp

- Asn Glu Asn Gln Arg Thr Pro Cys Thr Gly Asp Leu Met Val Arg Tyr
  755 760 765
- Val Gln Ala Gly Cys Leu Leu Pro Trp Phe Arg Asn His Tyr Asp Arg
  770 780
- Trp Ile Glu Ser Lys Asp His Gly Lys Asp Tyr Gln Glu Leu Tyr Met 785 790 795 800
- Tyr Pro Asn Glu Met Asp Thr Leu Arg Lys Phe Val Glu Phe Arg Tyr 805 810 815
- Arg Trp Gln Glu Val Leu Tyr Thr Ala Met Tyr Gln Asn Ala Ala Phe 820 825 830
- Gly Lys Pro Ile Ile Lys Ala Ala Ser Met Tyr Asn Asn Asp Ser Asn 835 840 845
- Val Arg Arg Ala Gln Asn Asp His Phe Leu Leu Gly Gly His Asp Gly 850 855 860
- Tyr Arg Ile Leu Cys Ala Pro Val Val Trp Glu Asn Ser Thr Glu Arg 865 870 875 880
- Glu Leu Tyr Leu Pro Val Leu Thr Gln Trp Tyr Lys Phe Gly Pro Asp 885 890 895
- Phe Asp Thr Lys Pro Leu Glu Gly Ala Met Asn Gly Gly Asp Arg Ile 900 905 910
- Tyr Asn Tyr Pro Val Pro Gln Ser Glu Ser Pro Ile Phe Val Arg Glu 915 920 925
- Gly Ala Ile Leu Pro Thr Arg Tyr Thr Leu Asn Gly Glu Asn Lys Ser 930 935 940
- Leu Asn Thr Tyr Thr Asp Glu Asp Pro Leu Val Phe Glu Val Phe Pro 945 950 955 960
- Leu Gly Asn Asn Arg Ala Asp Gly Met Cys Tyr Leu Asp Asp Gly Gly
  965 970 975
- Val Thr Thr Asn Ala Glu Asp Asn Gly Lys Phe Ser Val Val Lys Val
  980 985 990
- Ala Ala Glu Gln Asp Gly Gly Thr Glu Thr Ile Thr Phe Thr Asn Asp 995 1000 1005
- Cys Tyr Glu Tyr Val Phe Gly Gly Pro Phe Tyr Val Arg Val Arg 1010 1015 1020
- Gly Ala Gln Ser Pro Ser Asn Ile His Val Ser Ser Gly Ala Gly 1025 1030 1035
- Ser Gln Asp Met Lys Val Ser Ser Ala Thr Ser Arg Ala Ala Leu 1040 1045 1050

Phe Asn Asp Gly Glu Asn Gly Asp Phe Trp Val Asp Gln Glu Thr 1055 1060 1065

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Val Val Pro Ala Val Arg Leu Ala Val Arg Lys Ser Asn Arg Leu Asn 35 40 45

Val Ser Met Ser Ala Leu Phe Asp Lys Pro Thr Ala Val Thr Gly Gly 50 55 60

Lys Asp Asn Pro Asp Asn Ile Asn Tyr Thr Thr Tyr Asp Tyr Val Pro 65 70 75 80

Val Trp Arg Phe Asp Pro Leu Ser Asn Thr Asn Trp Phe Ala Ala Gly 85 90 95

Ser Ser Thr Pro Gly Asp Ile Asp Asp Trp Thr Ala Thr Met Asn Val 100 105 110

Asn Phe Asp Arg Ile Asp Asn Pro Ser Phe Thr Leu Glu Lys Pro Val 115 120 125

Gln Val Gln Val Thr Ser Tyr Lys Asn Asn Cys Phe Arg Val Arg Phe 130 135 140

Asn Pro Asp Gly Pro Ile Arg Asp Val Asp Arg Gly Pro Ile Leu Gln 145 150 155 160

Gln Gln Leu Asn Trp Ile Arg Lys Gln Glu Gln Ser Lys Gly Phe Asp 165 170 175

Pro Lys Met Gly Phe Thr Lys Glu Gly Phe Leu Lys Phe Glu Thr Lys
180 185 190

Asp Leu Asn Val Ile Ile Tyr Gly Asn Phe Lys Thr Arg Val Thr Arg

205

200

195

Lys Arg Asp Gly Lys Gly Ile Met Glu Asn Asn Glu Val Pro Ala Gly Ser Leu Gly Asn Lys Cys Arg Gly Leu Met Phe Val Asp Arg Leu Tyr 230 Gly Thr Ala Ile Ala Ser Val Asn Glu Asn Tyr Arg Asn Asp Pro Asp Arg Lys Glu Gly Phe Tyr Gly Ala Gly Glu Val Asn Cys Glu Phe Trp Asp Ser Glu Gln Asn Arg Asn Lys Tyr Ile Leu Glu Arg Thr Gly Ile 280 Ala Met Thr Asn Tyr Asn Tyr Asp Asn Tyr Asn Tyr Asn Gln Ser Asp 290 295 300 Leu Ile Ala Pro Gly Tyr Pro Ser Asp Pro Asn Phe Tyr Ile Pro Met 315 Tyr Phe Ala Ala Pro Trp Val Val Lys Gly Cys Ser Gly Asn Ser Asp Glu Gln Tyr Ser Tyr Gly Trp Phe Met Asp Asn Val Ser Gln Thr Tyr Met Asn Thr Gly Gly Thr Ser Trp Asn Cys Gly Glu Glu Asn Leu Ala Tyr Met Gly Ala Gln Cys Gly Pro Phe Asp Gln His Phe Val Tyr 370 375 Gly Asp Gly Asp Gly Leu Glu Asp Val Val Gln Ala Phe Ser Leu Leu 390 Gln Gly Lys Glu Phe Glu Asn Gln Val Leu Asn Lys Arg Ala Val Met 405 410 Pro Pro Lys Tyr Val Phe Gly Tyr Phe Gln Gly Val Phe Gly Ile Ala 420 Ser Leu Leu Arg Glu Gln Arg Pro Glu Gly Gly Asn Asn Ile Ser Val Gln Glu Ile Val Glu Gly Tyr Gln Ser Asn Asn Phe Pro Leu Glu Gly Leu Ala Val Asp Val Asp Met Gln Gln Asp Leu Arg Val Phe Thr Thr 465 Lys Ile Glu Phe Trp Thr Ala Asn Lys Val Gly Thr Gly Gly Asp Ser 490 Asn Asn Lys Ser Val Phe Glu Trp Ala His Asp Lys Gly Leu Val Cys

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Gln	Thr	Asn 515	Val	Thr	Cys	Phe	Leu 520	Arg	Asn	Asp	Asn	Gly 525	Gly	Ala	Asp
Tyr	Glu 530	Val	Asn	Gln	Thr	Leu 535	Arg	Glu	Lys	Gly	Leu 540	Tyr	Thr	Lys	Asn
Asp 545	Ser	Leu	Thr	Asn	Thr 550	Asn	Phe	Gly	Thr	Thr 555	Asn	Asp	Gly	Pro	Ser 560
Asp	Ala	Tyr	Ile	Gly 565	His	Leu	Asp	Tyr	Gly 570	Gly	Gly	Gly	Asn	Cys 575	Asp
Ala	Leu	Phe	Pro 580	Asp	Trp	Gly	Arg	Pro 585	Gly	Val	Ala	Glu	Trp 590	Trp	Gly
Asp	Asn	Tyr 595	Ser	Lys	Leu	Phe	Lys 600	Ile	Gly	Leu	Asp	Phe 605	Val	Trp	Gln
Asp	Met 610	Thr	Val	Pro	Ala	Met 615	Met	Pro	His	Lys	Val 620	Gly	Asp	Ala	Val
Asp 625	Thr	Arg	Ser	Pro	Tyr 630	Gly	Trp	Pro	Asn	Glu 635	Asn	Asp	Pro	Ser	Asn 640
Gly	Arg	Tyr	Asn	Trp 645	Lys	Ser	Tyr	His	Pro 650	Gln	Val	Leu	Val	Thr 655	Asp
Met	Arg	Tyr	Glu 660	Asn	His	Gly	Arg	Glu 665	Pro	Met	Phe	Thr	Gln 670	Arg	Asn
Met	His	Ala 675	Tyr	Thr	Leu	Cys	Glu 680	Ser	Thr	Arg	Lys	Glu 685	Gly	Ile	Val
Ala	Asn 690	Ala	Asp	Thr	Leu	Thr 695	Lys	Phe	Arg	Arg	Ser 700	Tyr	Ile	Ile	Ser
Arg 705	Gly	Gly	Tyr	Ile	Gly 710	Asn	Gln	His	Phe	Gly 715	Gly	Met	Trp	Val	Gly 720
Asp	Asn	Ser	Ser	Ser 725	Gln	Arg	Tyr	Leu	Gln 730	Met	Met	Ile	Ala	Asn 735	Ile
Val	Asn	Met	Asn 740	Met	Ser	Cys	Leu	Pro 745	Leu	Val	Gly	Ser	Asp 750	Ile	Gly
Gly	Phe	Thr 755	Ser	Tyr	Asp	Gly	Arg 760	Asn	Val	Cys	Pro	Gly 765	Asp	Leu	Met
Val	Arg 770	Phe	Val	Gln	Ala	Gly 775	Cys	Leu	Leu	Pro	Trp 780	Phe	Arg	Asn	His
Tyr 785	Gly	Arg	Leu	Val	Glu 790	Gly	Lys	Gln	Glu	Gly 795	Lys	Tyr	Tyr	Gln	Glu 800
Leu	Tvr	Met	Tvr	Lvs	Asp	Glu	Met	Ala	Thr	Leu	Ara	Lvs	Phe	Ile	Glu

805 810 815

Phe Arg Tyr Arg Trp Gln Glu Val Leu Tyr Thr Ala Met Tyr Gln Asn 820 825 830

- Ala Ala Phe Gly Lys Pro Ile Ile Lys Ala Ala Ser Met Tyr Asp Asn 835 840 845
- Asp Arg Asn Val Arg Gly Ala Gln Asp Asp His Phe Leu Leu Gly Gly 850 855 860
- His Asp Gly Tyr Arg Ile Leu Cys Ala Pro Val Val Trp Glu Asn Thr 865 870 875 880
- Thr Ser Arg Asp Leu Tyr Leu Pro Val Leu Thr Lys Trp Tyr Lys Phe 885 890 895
- Gly Pro Asp Tyr Asp Thr Lys Arg Leu Asp Ser Ala Leu Asp Gly Gly
  900 905 910
- Gln Met Ile Lys Asn Tyr Ser Val Pro Gln Ser Asp Ser Pro Ile Phe 915 920 925
- Val Arg Glu Gly Ala Ile Leu Pro Thr Arg Tyr Thr Leu Asp Gly Ser 930 935 940
- Asn Lys Ser Met Asn Thr Tyr Thr Asp Lys Asp Pro Leu Val Phe Glu 945 950 955 960
- Val Phe Pro Leu Gly Asn Asn Arg Ala Asp Gly Met Cys Tyr Leu Asp 965 970 975
- Asp Gly Gly Ile Thr Thr Asp Ala Glu Asp His Gly Lys Phe Ser Val 980 985 990
- Ile Asn Val Glu Ala Leu Arg Lys Gly Val Thr Thr Thr Ile Lys Phe 995 1000 1005
- Ala Tyr Asp Thr Tyr Gln Tyr Val Phe Asp Gly Pro Phe Tyr Val 1010 1015 1020
- Arg Ile Arg Asn Leu Thr Thr Ala Ser Lys Ile Asn Val Ser Ser 1025 1030 1035
- Gly Ala Gly Glu Glu Asp Met Thr Pro Thr Ser Ala Asn Ser Arg 1040 1045 1050
- Ala Ala Leu Phe Ser Asp Gly Gly Val Gly Glu Tyr Trp Ala Asp 1055 1060 1065
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<213> Unknown

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Asp Thr Thr Pro Pro Lys Ser Thr Lys Phe Pro Lys Asn Trp His Gly 35 40 45

Val Asn Leu Arg Phe Asp Asp Gly Thr Leu Gly Val Val Gln Phe Ile 50 55 60

Arg Pro Cys Val Trp Arg Val Arg Tyr Asp Pro Gly Phe Lys Thr Ser 65 70 75 80

Asp Glu Tyr Gly Asp Glu Asn Thr Arg Thr Ile Val Gln Asp Tyr Met 85 90 95

Ser Thr Leu Ser Asn Lys Leu Asp Thr Tyr Arg Gly Leu Thr Trp Glu 100 105 110

Thr Lys Cys Glu Asp Ser Gly Asp Phe Phe Thr Phe Ser Ser Lys Val 115 120 125

Thr Ala Val Glu Lys Ser Glu Arg Thr Arg Asn Lys Val Gly Asp Gly 130 135 140

Leu Arg Ile His Leu Trp Lys Ser Pro Phe Arg Ile Gln Val Val Arg 145 150 155 160

Thr Leu Thr Pro Leu Lys Asp Pro Tyr Pro Ile Pro Asn Val Ala Ala 165 170 175

Ala Glu Ala Arg Val Ser Asp Lys Val Val Trp Gln Thr Ser Pro Lys 180 185 190

Thr Phe Arg Lys Asn Leu His Pro Gln His Lys Met Leu Lys Asp Thr 195 200 205

Val Leu Asp Ile Val Lys Pro Gly His Gly Glu Tyr Val Gly Trp Gly 210 215 220

Glu Met Gly Gly Ile Gln Phe Met Lys Glu Pro Thr Phe Met Asn Tyr 225 230 235 240

Phe Asn Phe Asp Asn Met Gln Tyr Gln Gln Val Tyr Ala Gln Gly Ala 245 250 255

- Leu Asp Ser Arg Glu Pro Leu Tyr His Ser Asp Pro Phe Tyr Leu Asp 260 265 270
- Val Asn Ser Asn Pro Glu His Lys Asn Ile Thr Ala Thr Phe Ile Asp 275 280 285
- Asn Tyr Ser Gln Ile Ala Ile Asp Phe Gly Lys Thr Asn Ser Gly Tyr 290 295 300
- Ile Lys Leu Gly Thr Arg Tyr Gly Gly Ile Asp Cys Tyr Gly Ile Ser 305 310 315 320
- Ala Asp Thr Val Pro Glu Ile Val Arg Leu Tyr Thr Gly Leu Val Gly 325 330 335
- Arg Ser Lys Leu Lys Pro Arg Tyr Ile Leu Gly Ala His Gln Ala Cys 340 345 350
- Tyr Gly Tyr Gln Gln Glu Ser Asp Leu Tyr Ser Val Val Gln Gln Tyr 355 360 365
- Arg Asp Cys Lys Phe Pro Leu Asp Gly Ile His Val Asp Val Asp Val 370 380
- Gln Asp Gly Phe Arg Thr Phe Thr Thr Asn Pro His Thr Phe Pro Asn 385 390 395 400
- Pro Lys Glu Met Phe Thr Asn Leu Arg Asn Asn Gly Ile Lys Cys Ser 405 410 415
- Thr Asn Ile Thr Pro Val Ile Ser Ile Asn Asn Arg Glu Gly Gly Tyr 420 425 430
- Ser Thr Leu Leu Glu Gly Val Asp Lys Lys Tyr Phe Ile Met Asp Asp 435 440 445
- Arg Tyr Thr Glu Gly Thr Ser Gly Asn Ala Lys Asp Val Arg Tyr Met 450 455 460
- Tyr Tyr Gly Gly Asn Lys Val Glu Val Asp Pro Asn Asp Val Asn 465 470 475 480
- Gly Arg Pro Asp Phe Lys Asp Asn Tyr Asp Phe Pro Ala Asn Phe Asn 485 490 495
- Ser Lys Gln Tyr Pro Tyr His Gly Gly Val Ser Tyr Gly Tyr Gly Asn 500 505 510
- Gly Ser Ala Gly Phe Tyr Pro Asp Leu Asn Arg Lys Glu Val Arg Ile 515 520 525
- Trp Trp Gly Met Gln Tyr Lys Tyr Leu Phe Asp Met Gly Leu Glu Phe 530 540
- Val Trp Gln Asp Met Thr Thr Pro Ala Ile His Thr Ser Tyr Gly Asp 545 550 555 560

- Met Lys Gly Leu Pro Thr Arg Leu Leu Val Thr Ser Asp Ser Val Thr 565 570 575
- Asn Ala Ser Glu Lys Lys Leu Ala Ile Glu Thr Trp Ala Leu Tyr Ser 580 585 590
- Tyr Asn Leu His Lys Ala Thr Trp His Gly Leu Ser Arg Leu Glu Ser 595 600 605
- Arg Lys Asn Lys Arg Asn Phe Ile Leu Gly Arg Gly Ser Tyr Ala Gly 610 615 620
- Ala Tyr Arg Phe Ala Gly Leu Trp Thr Gly Asp Asn Ala Ser Asn Trp 625 630 635 640
- Glu Phe Trp Lys Ile Ser Val Ser Gln Val Leu Ser Leu Gly Leu Asn 645 650 655
- Gly Val Cys Ile Ala Gly Ser Asp Thr Gly Gly Phe Glu Pro Tyr Arg
  660 665 670
- Asp Ala Asn Gly Val Glu Glu Lys Tyr Cys Ser Pro Glu Leu Leu Ile 675 680 685
- Arg Trp Tyr Thr Gly Ser Phe Leu Leu Pro Trp Leu Arg Asn His Tyr 690 695 700
- Val Lys Lys Asp Arg Lys Trp Phe Gln Glu Pro Tyr Ser Tyr Pro Lys 705 710 715 720
- His Leu Glu Thr His Pro Glu Leu Ala Asp Gln Ala Trp Leu Tyr Lys
  725 730 735
- Ser Val Leu Glu Ile Cys Arg Tyr Tyr Val Glu Leu Arg Tyr Ser Leu 740 745 750
- Ile Gln Leu Leu Tyr Asp Cys Met Phe Gln Asn Val Val Asp Gly Met 755 760 765
- Pro Ile Thr Arg Ser Met Leu Leu Thr Asp Thr Glu Asp Thr Thr Phe
  770 780
- Phe Asn Glu Ser Gln Lys Phe Leu Asp Asn Gln Tyr Met Ala Gly Asp 785 790 795 800
- Asp Ile Leu Val Ala Pro Ile Leu His Ser Arg Lys Glu Ile Pro Gly 805 810 815
- Glu Asn Arg Asp Val Tyr Leu Pro Leu Tyr His Thr Trp Tyr Pro Ser 820 825 830
- Asn Leu Arg Pro Trp Asp Asp Gln Gly Val Ala Leu Gly Asn Pro Val 835 840 845
- Glu Gly Gly Ser Val Ile Asn Tyr Thr Ala Arg Ile Val Ala Pro Glu 850 855 860

Asp Tyr Asn Leu Phe His Ser Val Val Pro Val Tyr Val Arg Glu Gly 865 870 875 880

Ala Ile Ile Pro Gln Ile Glu Val Arg Gln Trp Thr Gly Gln Gly Gly 885 890 895

Ala Asn Arg Ile Lys Phe Asn Ile Tyr Pro Gly Lys Asp Lys Glu Tyr 900 905 910

Cys Thr Tyr Leu Asp Asp Gly Val Ser Arg Asp Ser Ala Pro Glu Asp 915 920 925

Leu Pro Gln Tyr Lys Glu Thr His Glu Gln Ser Lys Val Glu Gly Ala 930 935 940

Glu Ile Ala Lys Gln Ile Gly Lys Lys Thr Gly Tyr Asn Ile Ser Gly 945 950 955 960

Thr Asp Pro Glu Ala Lys Gly Tyr His Arg Lys Val Ala Val Thr Gln 965 970 975

Thr Ser Lys Asp Lys Thr Arg Thr Val Thr Ile Glu Pro Lys His Asn 980 985 990

Gly Tyr Asp Pro Ser Lys Glu Val Gly Asp Tyr Tyr Thr Ile Ile Leu 995 1000 1005

Trp Tyr Ala Pro Gly Phe Asp Gly Ser Ile Val Asp Val Ser Lys 1010 1015 1020

Thr Thr Val Asn Val Glu Gly Gly Val Glu His Gln Val Tyr Lys 1025 1030 1035

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<212> PRT

<213> Unknown

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<223> fungus sp. or fungus infected gracilariopsis sp.

<400> 4

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Asp Gln Thr Pro Pro Gln Gly Thr Lys Asp Pro Lys Ser Trp His Ala 35 40 45

Val Asn Leu Pro Phe Asp Asp Gly Thr Met Cys Val Val Gln Phe Val Arg Pro Cys Val Trp Arg Val Arg Tyr Asp Pro Ser Val Lys Thr Ser 70 Asp Glu Tyr Gly Asp Glu Asn Thr Arg Thr Ile Val Gln Asp Tyr Met Thr Thr Leu Val Gly Asn Leu Asp Ile Phe Arg Gly Leu Thr Trp Val 105 Ser Thr Leu Glu Asp Ser Gly Glu Tyr Tyr Thr Phe Lys Ser Glu Val 115 Thr Ala Val Asp Glu Thr Glu Arg Thr Arg Asn Lys Val Gly Asp Gly Leu Lys Ile Tyr Leu Trp Lys Asn Pro Phe Arg Ile Gln Val Val Arg 150 155 Leu Leu Thr Pro Leu Val Asp Pro Phe Pro Ile Pro Asn Val Ala Asn 170 Ala Thr Ala Arg Val Ala Asp Lys Val Val Trp Gln Thr Ser Pro Lys 185 Thr Phe Arg Lys Asn Leu His Pro Gln His Lys Met Leu Lys Asp Thr 195 Val Leu Asp Ile Ile Lys Pro Gly His Gly Glu Tyr Val Gly Trp Gly Glu Met Gly Gly Ile Glu Phe Met Lys Glu Pro Thr Phe Met Asn Tyr Phe Asn Phe Asp Asn Met Gln Tyr Gln Gln Val Tyr Ala Gln Gly Ala Leu Asp Ser Arg Glu Pro Leu Tyr His Ser Asp Pro Phe Tyr Leu Asp 265 Val Asn Ser Asn Pro Glu His Lys Asn Ile Thr Ala Thr Phe Ile Asp 275 Asn Tyr Ser Gln Ile Ala Ile Asp Phe Gly Lys Thr Asn Ser Gly Tyr 295 Ile Lys Leu Gly Thr Arg Tyr Gly Gly Ile Asp Cys Tyr Gly Ile Ser Ala Asp Thr Val Pro Glu Ile Val Arg Leu Tyr Thr Gly Leu Val Gly 325 Arg Ser Lys Leu Lys Pro Arg Tyr Ile Leu Gly Ala His Gln Ala Cys

345

340

Tyr Gly Tyr Gln Gln Glu Ser Asp Leu His Ala Val Val Gln Gln Tyr Arg Asp Thr Lys Phe Pro Leu Asp Gly Leu His Val Asp Val Asp Phe 370 375 380 Gln Asp Asn Phe Arg Thr Phe Thr Thr Asn Pro Ile Thr Phe Pro Asn 390 395 Pro Lys Glu Met Phe Thr Asn Leu Arg Asn Asn Gly Ile Lys Cys Ser 410 Thr Asn Ile Thr Pro Val Ile Ser Ile Arg Asp Arg Pro Asn Gly Tyr 420 Ser Thr Leu Asn Glu Gly Tyr Asp Lys Lys Tyr Phe Ile Met Asp Asp Arg Tyr Thr Glu Gly Thr Ser Gly Asp Pro Gln Asn Val Arg Tyr Ser 450 455 Phe Tyr Gly Gly Asn Pro Val Glu Val Asn Pro Asn Asp Val Trp Ala Arg Pro Asp Phe Gly Asp Asn Tyr Asp Phe Pro Thr Asn Phe Asn Cys Lys Asp Tyr Pro Tyr His Gly Gly Val Ser Tyr Gly Tyr Gly Asn 505 Gly Thr Pro Gly Tyr Tyr Pro Asp Leu Asn Arg Glu Glu Val Arg Ile 520 Trp Trp Gly Leu Gln Tyr Glu Tyr Leu Phe Asn Met Gly Leu Glu Phe 535 Val Trp Gln Asp Met Thr Thr Pro Ala Ile His Ser Ser Tyr Gly Asp 545 550 555 Met Lys Gly Leu Pro Thr Arg Leu Leu Val Thr Ala Asp Ser Val Thr 570 Asn Ala Ser Glu Lys Lys Leu Ala Ile Glu Ser Trp Ala Leu Tyr Ser Tyr Asn Leu His Lys Ala Thr Phe His Gly Leu Gly Arg Leu Glu Ser 595 Arg Lys Asn Lys Arg Asn Phe Ile Leu Gly Arg Gly Ser Tyr Ala Gly Ala Tyr Arg Phe Ala Gly Leu Trp Thr Gly Asp Asn Ala Ser Thr Trp 625 630 Glu Phe Trp Lys Ile Ser Val Ser Gln Val Leu Ser Leu Gly Leu Asn

650

645

Gly Val Cys Ile Ala Gly Ser Asp Thr Gly Gly Phe Glu Pro Ala Arg 665 Thr Glu Ile Gly Glu Glu Lys Tyr Cys Ser Pro Glu Leu Leu Ile Arg 675 680 Trp Tyr Thr Gly Ser Phe Leu Leu Pro Trp Leu Arg Asn His Tyr Val 695 Lys Lys Asp Arg Lys Trp Phe Gln Glu Pro Tyr Ala Tyr Pro Lys His 710 715 Leu Glu Thr His Pro Glu Leu Ala Asp Gln Ala Trp Leu Tyr Lys Ser 725 Val Leu Glu Ile Cys Arg Tyr Trp Val Glu Leu Arg Tyr Ser Leu Ile 745 Gln Leu Leu Tyr Asp Cys Met Phe Gln Asn Val Val Asp Gly Met Pro Leu Ala Arg Ser Met Leu Leu Thr Asp Thr Glu Asp Thr Thr Phe Phe Asn Glu Ser Gln Lys Phe Leu Asp Asn Gln Tyr Met Ala Gly Asp Asp 790 Ile Leu Val Ala Pro Ile Leu His Ser Arg Asn Glu Val Pro Gly Glu 805 Asn Arg Asp Val Tyr Leu Pro Leu Phe His Thr Trp Tyr Pro Ser Asn 825 Leu Arg Pro Trp Asp Asp Gln Gly Val Ala Leu Gly Asn Pro Val Glu Gly Gly Ser Val Ile Asn Tyr Thr Ala Arg Ile Val Ala Pro Glu Asp 855 Tyr Asn Leu Phe His Asn Val Val Pro Val Tyr Ile Arg Glu Gly Ala Ile Ile Pro Gln Ile Gln Val Arg Gln Trp Ile Gly Glu Gly Gly Pro Asn Pro Ile Lys Phe Asn Ile Tyr Pro Gly Lys Asp Lys Glu Tyr Val 900 905 Thr Tyr Leu Asp Asp Gly Val Ser Arg Asp Ser Ala Pro Asp Asp Leu 920 Pro Gln Tyr Arg Glu Ala Tyr Glu Gln Ala Lys Val Glu Gly Lys Asp 930 935 Val Gln Lys Gln Leu Ala Val Ile Gln Gly Asn Lys Thr Asn Asp Phe 945 950 955

Ser Ala Ser Gly Ile Asp Lys Glu Ala Lys Gly Tyr His Arg Lys Val 965 970 975

Ser Ile Lys Gln Glu Ser Lys Asp Lys Thr Arg Thr Val Thr Ile Glu 980 985 990

Pro Lys His Asn Gly Tyr Asp Pro Ser Lys Glu Val Gly Asn Tyr Tyr 995 1000 1005

Thr Ile Ile Leu Trp Tyr Ala Pro Gly Phe Asp Gly Ser Ile Val 1010 1015 1020

Asp Val Ser Gln Ala Thr Val Asn Ile Glu Gly Gly Val Glu Cys 1025 1030 1035

Glu Ile Phe Lys Asn Thr Gly Leu His Thr Val Val Val Asn Val 1040 1045 1050

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Thr Ala 1070

<210> 5

<211> 1092

<212> PRT

<213> Unknown

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<223> fungus sp. or fungus infected gracilariopsis sp.

<400> 5

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Thr Phe Val Gly Ala Asp Ile Arg Ser Gly Ile Arg Ile Gln Ser Ala 20 25 30

Leu Pro Ala Val Arg Asn Ala Val Arg Arg Ser Lys His Tyr Asn Val
35 40 45

Ser Met Thr Ala Leu Ser Asp Lys Gln Thr Ala Ile Ser Ile Gly Pro 50 55 60

Asp Asn Pro Asp Gly Ile Asn Tyr Gln Asn Tyr Asp Tyr Ile Pro Val 65 70 75 80

Ala Gly Phe Thr Pro Leu Ser Asn Thr Asn Trp Tyr Ala Ala Gly Ser 85 90 95

Ser Thr Pro Gly Gly Ile Thr Asp Trp Thr Ala Thr Met Asn Val Lys
100 105 110

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120

115

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170

175

165

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<210> 10

<211> 3213

<212> DNA

<213> Unknown

<220>

<223> fungus sp. or fungus infected gracilariopsis sp.

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<220>

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<sup>&</sup>lt;211> 3279

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Unknown

<sup>&</sup>lt;223> fungus sp. or fungus infected gracilariopsis sp.

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<sup>&</sup>lt;210> 12

<sup>&</sup>lt;211> 1712

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Unknown

<sup>&</sup>lt;220>

<sup>&</sup>lt;223> fungus sp. or fungus infected gracilariopsis sp.

<sup>&</sup>lt;400> 12

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